1 ABSTRACT

2	The present invention provides a method of identifying specific over-
3	expressed lung cancer genes using a novel in-situ screening approach and further relates
4	to a composition comprising novel genes that are identified using the present method.
5	The present invention further contemplates providing a high-throughput molecular
6	classification and diagnostics methodology based on detecting mRNAs for the identified
7	lung-cancer-overexpressed genes.
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